|       | 1      | 1        | :      |       |
|-------|--------|----------|--------|-------|
| EKUT. | a squa | lene epo | xidase | gene. |
| ,     |        | P -      |        | G,    |

or

ii) t-HMG, an HMG-Co-A-reductase gene, and ERG9, a squalene synthetase gene,

or

t-HMG, an HMG-Co-A-reductase gene, andSAT1, an acyl-CoA: sterol-acyl transferase gene,

or

iv) t-HMG, an HMG-Co-A-reductase gene, and ERG1, a squalene epoxidase gene,

or

v) ERG9, a squalene synthetase gene, and SAT1, an acyl-CoA: sterol-acyl transferase gene,

or

vi) ERG9, a squalene synthetase gene, and ERG1, a squalene epoxidase gene,

or

vii) SAT1, an acyl-CoA: sterol-acyl transferase gene, and ERG1, a squalene epoxidase gene,

or

- viii) one of the genes selected from the group consisting of ERG9, SAT1 and ERG1,
- b) transforming a microorganism with a plasmid mentioned in i) to vii), or, simultaneously or in succession, with two or more of the plasmids mentioned in viii), and
- c) culturing the transformed microorganism under conditions in which it produces ergosterol and an intermediate product of ergosterol biosythesis.
- 36. (Amended) A yeast strain S. cerevisiae AH22 comprising at least one gene selected from the group consisting of t-HMG, an HMG-Co-A-reductase gene, ERG9, a squalene synthetase gene; SAT1, an Acyl-CoA sterol-acyl transferase gene; and ERG1, a squalene epoxidase gene.
  - 37. (Amended) The plasmid YEpH2, which comprises the ADH-promoter, the t-HMG

gene, and the TRP-terminator, as shown in Fig. 1.

- 38. (Amended) The plasmid YDpUHK3, which comprises the ADH-promoter, the t-HMG gene, the TRP-terminator, the gene for kanamycin resistance and the ura3 gene, as shown in Fig. 2.
- 39. (Amended) The plasmid pADL-SAT1, which comprises the SAT1 gene and the LEU2 gene of YEp13, as shown in Fig 3.
- 42. (Amended) A method for producing an intermediate sterol product with a 5,7-diene structure in the biosynthesis of ergosterol, comprising transforming a microorganism with a plasmid according to claim 37, and culturing the transformed microorganism under conditions in which it produces said intermediate sterol product.
- 43. (Amended) An expression cassette that comprises a t-HMG gene operatively linked to an ADH-promoter and a TRP-terminator, and an SAT1 gene operatively linked to an ADH-promoter and a TRP-terminator.
- 44. (Amended) An expression cassette that comprises a t-HMG gene operatively linked to an ADH-promoter and a TRP-terminator, and an SAT1 gene operatively linked to an ADH-promoter and a TRP-terminator, and an ERG9-gene operatively linked to an ADH-promoter and a TRP-terminator.
- 53. (Amended) A method for producing ergosterol or one or more intermediate products of its biosynthesis, comprising expressing in a microorganism a plasmid which comprises the following genes:
  - t-HMG, an HMG-Co-A-reductase gene,
    ERG9, a squalene synthetase gene,
    SAT1, an Acyl-CoA: sterol-acyl transferase gene, and
    ERG1, a squalene epoxidase gene,

or

ii) t-HMG, an HMG-Co-A-reductase gene, and

|       |          | ERG9, a squalene synthetase gene,  |
|-------|----------|--|
|       | or       |  |
|       | iii)     | t-HMG, an HMG-Co-A-reductase gene, and                                     |
|       |          | SAT1, an acyl-CoA: sterol-acyl transferase gene,                           |
|       | or       |  |
|       | iv)      | t-HMG, an HMG-Co-A-reductase gene, and                                     |
|       |          | ERG1, a squalene epoxidase gene,   |
|       | or       |  |
|       | v)       | ERG9, a squalene synthetase gene, and                                      |
|       |          | SAT1, an acyl-CoA: sterol-acyl transferase gene,                           |
|       | or       |  |
|       | vi)      | ERG9, a squalene synthetase gene, and                                      |
|       |          | ERG1, a squalene epoxidase gene,   |
|       | or       |  |
|       | vii)     | SAT1, an acyl-CoA: sterol-acyl transferase gene, and                       |
|       |          | ERG1, a squalene epoxidase gene,   |
|       | or       |  |
|       | viii)    | one of the genes selected from the group consisting of ERG9, SAT1 and ERG1 |
| and i | solating | the expressed ergosterol or intermediate products of its biosynthesis.     |
|       |          |  |